PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/890,475

DATE: 01/22/2002
TIME: 16:34:40

Input Set : **A:\mewburn.app**
Output Set: N:\CRF3\01182002\I890475.raw

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 3 <110> APPLICANT: Johanson, Urban
 4         West, Joanne
 5         Dean, Caroline
 7 <120> TITLE OF INVENTION: Arabidopsis thaliana derived Frigida gene conferring late
flowering
 9 <130> FILE REFERENCE: Mewburn
11 <140> CURRENT APPLICATION NUMBER: US 09/890,475
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C--> 12 **<141> CURRENT FILING DATE: 2001-11-13**
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14 <150> PRIOR APPLICATION NUMBER: PCT/GB00/00197
15 <151> PRIOR FILING DATE: 2000-01-25
17 <150> PRIOR APPLICATION NUMBER: GB 9902660.1
18 <151> PRIOR FILING DATE: 1999-02-05
20 <160> NUMBER OF SEQ ID NOS: 58
22 <170> SOFTWARE: PatentIn Ver. 2.1
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25 <211> LENGTH: 609
26 <212> TYPE: PRT
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Description of Artificial Sequence: FRI amino acid
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38              20                  25                  30
40 Pro Lys Ile Val Glu Thr Glu Ser Thr Ser Met Asp Ile Thr Ile Gly
41          35                  40                  45
43 Gln Ser Lys Gln Pro Gln Phe Leu Lys Ser Ile Asp Glu Leu Ala Ala
44       50                  55                  60
46 Phe Ser Val Ala Val Glu Thr Phe Lys Arg Gln Phe Asp Asp Leu Gln
47  65                  70                  75                  80
49 Lys His Ile Glu Ser Ile Glu Asn Ala Ile Asp Ser Lys Leu Glu Ser
50                  85                  90                  95
52 Asn Gly Val Val Leu Ala Ala Arg Asn Asn Asn Phe His Gln Pro Met
53              100                 105                 110
55 Leu Ser Pro Pro Arg Asn Asn Val Ser Val Glu Thr Thr Val Thr Val
56          115                 120                 125
58 Ser Gln Pro Ser Gln Glu Ile Val Pro Glu Thr Ser Asn Lys Pro Glu
59       130                 135                 140
61 Gly Gly Arg Met Cys Glu Leu Met Cys Ser Lys Gly Leu Arg Lys Tyr
62 145                 150                 155                 160
64 Ile Tyr Ala Asn Ile Ser Asp Gln Ala Lys Leu Met Glu Glu Ile Pro
65                 165                 170                 175
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73 Pro Met Ser Ser Ala Arg Gln Val Ser Leu Leu Ile Leu Glu Ser Phe
74         210                 215                 220
76 Leu Leu Met Pro Asp Arg Gly Lys Gly Lys Val Lys Ile Glu Ser Trp
77 225                 230                 235                 240
79 Ile Lys Asp Glu Ala Glu Thr Ala Ala Val Ala Trp Arg Lys Arg Leu
80             245                 250                 255
82 Met Thr Glu Gly Gly Leu Ala Ala Ala Glu Lys Met Asp Ala Arg Gly
83             260                 265                 270
85 Leu Leu Leu Leu Val Ala Cys Phe Gly Val Pro Ser Asn Phe Arg Ser
86         275                 280                 285
88 Thr Asp Leu Leu Asp Leu Ile Arg Met Ser Gly Ser Asn Glu Ile Ala
89     290                 295                 300
91 Gly Ala Leu Lys Arg Ser Gln Phe Leu Val Pro Met Val Ser Gly Ile
92 305                 310                 315                 320
94 Val Glu Ser Ser Ile Lys Arg Gly Met His Ile Glu Ala Leu Glu Met
95             325                 330                 335
97 Val Tyr Thr Phe Gly Met Glu Asp Lys Phe Ser Ala Ala Leu Val Leu
98             340                 345                 350
100 Thr Ser Phe Leu Lys Met Ser Lys Glu Ser Phe Glu Arg Ala Lys Arg
101         355                 360                 365
103 Lys Ala Gln Ser Pro Leu Ala Phe Lys Glu Ala Ala Thr Lys Gln Leu
104     370                 375                 380
106 Ala Val Leu Ser Ser Val Met Gln Cys Met Glu Thr His Lys Leu Asp
107 385                 390                 395                 400
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110             405                 410                 415
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127 Tyr Leu Gly Pro Ser Thr Ser Phe Pro His Arg Ser Arg Arg Ser Pro
128         500                 505                 510
130 Glu Tyr Met Val Pro Leu Pro His Gly Gly Leu Gly Arg Ser Val Tyr
131         515                 520                 525
133 Ala Tyr Glu His Leu Ala Pro Asn Ser Tyr Ser Pro Gly His Gly His
134     530                 535                 540
136 Arg Leu His Arg Gln Tyr Ser Pro Ser Leu Val His Gly Gln Arg His
137 545                 550                 555                 560
139 Pro Leu Gln Tyr Ser Pro Pro Ile His Gly Gln Gln Gln Leu Pro Tyr
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142 Gly Ile Gln Arg Val Tyr Arg His Ser Pro Ser Glu Glu Arg Tyr Leu
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148 Lys
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154 <211> LENGTH: 3761
155 <212> TYPE: DNA
156 <213> ORGANISM: Arabidopsis thaliana
158 <400> SEQUENCE: 2
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161 cttttgggtt catattaccg agcaagaacc gttatttgtg attagacatg ttataaacca 180
162 ctgctttagt gactatttaa aacaatatat tacatgtcgt aatcatgcaa cctaactatg 240
163 ttttcattaa tcaaatacaa agaataaaga gaaaagtgcg tagattcaat tatttggcat 300
164 agactcaaaa gagtgtatat atatctgact tttattaaat tattaaacac aaatacatat 360
165 tttcataagc aaaactataa aagccctaaa catataatga ttacctcaaa ggaaaaagtc 420
166 gttttctcct acttaaaaga taggttactt cctaattaat atataattta tgtgaacttc 480
167 acaatataca gttcaataaa atttggtaat ttgaccgatt taaggagagt ggaaattagg 540
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169 gcgcaaccca caacgacggc gaatccactg ctgcagcgac atcaatctga acagcgacga 660
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172 gaaacattca aacgccaatt cgatgatctt cagaagcaca tcgagtcaat cgaaaacgca 840
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230 <223> OTHER INFORMATION: Description of Artificial Sequence: likely cDNA
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236 tattaaatta ttaaacacaa atacatattt tcataagcaa aactataaaa gccctaaaca 180
237 tataatgatt acctcaaagg aaaaagtcgt tttctcctac ttaaaagata ggttacttcc 240
238 taattaatat ataatttatg tgaacttcac aatatacagt tcaataaaat ttggtaattt 300
239 gaccgattta aggagagtgg aaattagggc ttctgcaatc ttttttcttc gccgcaatct 360
240 catgtccaat tatccaccga cggtggcggc gcaacccaca acgacggcga tccactgct 420
241 gcagcgacat caatctgaac agcgacgaag agaattaccg aagattgtcg aaacagagtc 480
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247 gaataaaccg gagggggggac gtatgtgtga gttgatgtgt agcaaaggtc tgcgtaaata 840
248 catatacgcg aatatctctg atcaagctaa gttaatggaa gagattcctt cagctttgaa 900
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253 aggattagct gcggctgaga aaatggatgc aagggggtttg cttttactag ttgcttgttt 1200
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260 tcctgcgaaa gaactaccag gatggcagat caaagagcaa attgttagct tggagaaaga 1620
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263 ggaaatgcca ccagtaactt cttcatcgta ttctcctatc taccgtgata gaagctttcc 1800
264 tagtcaaaga gacgatgacc aagatgaaat atcagctctt gtgagtagtt acctcggccc 1860
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270 tcgcagtaac tcatcattag accccaaata ggaggaatgt aaatttgtaa caaagctttt 2220
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276 <212> TYPE: DNA
277 <213> ORGANISM: Artificial Sequence
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280 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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VERIFICATION SUMMARY   DATE: 01/22/2002
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